FILE 'CAPLUS' ENTERED AT 15:55:19 ON 02 SEP 1999

- key terms

L1 2901 SEA ABB=ON PLU=ON PNEUMOLYS? OR PNEUMOCOC?

L2 95 SEA ABB=ON PLU=ON L1 AND MODIF?

L3 26 SEA ABB=ON PLU=ON L2 AND (MUTAT? OR MUTANT OR MUTAGEN? OR POLYMORPH?)

=> d 1-26 .beverly

L3 ANSWER 1 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1999:187746 CAPLUS

DOCUMENT NUMBER:

131:29594

TITLE:

Penicillin-binding protein-mediated resistance

in pneumococci and staphylococci

AUTHOR (S):

Chambers, Henry F.

CORPORATE SOURCE:

Medical Service, San Francisco General Hospital,

and Department of Medicine, University of California, San Francisco, CA, 94143, USA

SOURCE:

J. Infect. Dis. (1999), 179(Suppl. 2), S353-S359

CODEN: JIDIAQ; ISSN: 0022-1899

PUBLISHER:

University of Chicago Press

DOCUMENT TYPE:

Journal; General Review

LANGUAGE:

English

A review with 58 refs. Target alteration underlies resistance to AB .beta.-lactam antibiotics in both Staphylococcus species and Streptococcus pneumoniae. The penicillin-binding protein (PBP) targets in penicillin-resistant strains of S. pneumoniae are modified, low-binding-affinity versions of the native PBPs. Multiple PBP targets may be modified by transformation and homologous recombination with DNA from PBP genes of viridans streptococci. The level of resistance is detd. by how many and to what extent targets are modified. In contrast, methicillin resistance in staphylococci is due to expression of PBP 2a, a novel, low-affinity PBP for which there is no homolog in methicillin-susceptible strains. PBP 2a is encoded by mecA, a highly conserved gene most likely acquired by a rare transposition from Staphylococcus sciuri or a closely related ancestor. Expression of resistance can be highly variable, but this seems not to be detd. by PBP modifications. Several non-PBP factors are required for high-level resistance.

L3 ANSWER 2 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1999:77590 CAPLUS

DOCUMENT NUMBER:

130:152551

TITLE:

Modified immunogenic

pneumolysin compositions as vaccines

INVENTOR (S):

Minetti, Conceicao; Michon, Francis; Pullen, Jeffrey K.; Polvino-Bodnar, Maryellen; Liang,

Shu-Mei; Tai, Joseph Y.

PATENT ASSIGNEE(S):

North American Vaccine, Inc., USA

Searcher :

Shears 308-4994

SOURCE:

PCT Int. Appl., 116 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

	PA'	PATENT NO.				KIND		DATE			APPLICATION NO.					DATE		
										-								
	WO	9903	884		Α	2	1999	0128		W	0 19	98-U	S147	16	1998	0721		
	WO	9903884			A3		19990408			WO 1998-US14716 19980721								
		W:	AL,	AM,	ΑT,	AU,	ΑZ,	BA,	BB,	BG,	BR,	BY.	CA.	CH.	CN.	CU.	CZ.	
			DE,	DK,	EE,	ES,	FI,	GB,	GE,	GH,	GM,	HR.	HU.	ID.	TI	TS,	.TD	
			KE,	KG,	KP,	KR,	KZ,	LC,	LK,	LR,	LS.	LT.	LU.	LV.	MD.	MG,	MK	
			MN,	MW,	MX,	NO,	NZ,	PL,	PT,	RO,	RU.	SD.	SE.	SG.	SI.	SK	SI.	
			TJ,	TM,	TR,	TT,	UA,	UG,	US,	US,	UZ,	VN.	YU.	ZW.	ΔM.	Δ7.	BV	
			KG,	KZ,	MD,	RU,	TJ,	TM	•	•	•		,	,	,	,	<i>D</i> 1,	
		RW:	GH,	GM,	KE,	LS,	MW,	SD,	SZ,	UG,	ZW,	AT,	BE.	CH.	CY.	DE.	DK	
			ES,	FI,	FR,	GB,	GR,	IE,	IT,	LU,	MC,	NL.	PT.	SE.	BF.	B.T.	CE.	
			CG,	CI,	CM,	GA,	GN,	GW,	ML,	MR,	NE,	SN.	TD.	TG	,	20,	Q. ,	
AU 9884078 A1							19990	210		AU 1998-84078 19980721								
PRIO		APPI								US 1997-5330								
										US 1998-73456 19980202								
										US 1998-345697								
															19980			

AΒ This invention relates to modified pneumolysin polypeptides that retain the immunogenic nature of pneumolysin but have reduced or undetectable hemolytic activity compared to native pneumolysin. The invention also provides a method for generating novel pneumolysin variants with these desired characteristic properties. The invention also provides immunogenic compns. useful as pharmaceutical compns. including vaccines in which non-toxic, modified pneumolysin is used to stimulate protective immunity against Streptococcus pneumoniae. The vaccines may be compns. in which the modified pneumolysin in conjugated to bacterial polysaccharides or may be carried on an attenuated viral vector. In addn., the invention also provides a method of using the non-toxic, modified pneumolysin toxoid in order to stimulate antibodies against Streptococcus pneumoniae in a treated individual which are then isolated and transferred to a second individual, thereby conferring protection against Streptococcus pneumoniae in the second individual. Prepd. and tested for immunogenicity were polypeptides pNVJ1, pNVJ20, pNVJ22, pNVJ45, pNVJ56, pNVJ103, pNVJ207, pNVJ111, and pNVJ211 and corresponding nucleic acid sequences.

L3 ANSWER 3 OF 26 CAPLUS COPYRIGHT 1999 ACS ACCESSION NUMBER: 1998:783992 CAPLUS

DOCUMENT NUMBER:

130:120724

TITLE:

The molecular mechanism of pneumolysin , a virulence factor from Streptococcus

pneumoniae

AUTHOR (S):

Rossjohn, Jamie; Gilbert, Robert J. C.; Crane, Dennis; Morgan, Peter J.; Mitchell, Timothy J.; Rowe, Arthur J.; Andrew, Peter W.; Paton, James

C.; Tweten, Rodney K.; Parker, Michael W.

CORPORATE SOURCE:

The Ian Potter Foundation Protein

Crystallography Laboratory, St. Vincent's Institute of Medical Research, Fitzroy, 3065,

Australia

SOURCE:

J. Mol. Biol. (1998), 284(2), 449-461

CODEN: JMOBAK; ISSN: 0022-2836

PUBLISHER:

Academic Press

DOCUMENT TYPE:

Journal English

LANGUAGE:

Pneumolysin, a member of the thiol-activated cytolysin family of toxins, is a virulence factor from the Gram-pos. bacterium Streptococcus pneumoniae. The toxin forms large oligomeric pores in cholesterol-contg. membranes of eukaryotic cells. A plethora of biochem. and mutagenesis data have been published on pneumolysin, since its initial characterization in the 1930s. Here we present an homol. model of the monomeric and oligomeric forms of pneumolysin based on the recently detd. crystal structure of perfringolysin O and electron microscopy data. A feature of the model is a striking electroneg. surface on parts of pneumolysin that may reflect its cytosolic location in the bacterial cell. The models provide a mol. basis for understanding the effects of published mutagenesis and biochem. modifications on the toxic activity of pneumolysin. In addn., spectroscopic data are presented that shed new light on pneumolysin activity and have guided us to hypothesize a detailed model of membrane insertion. These data show that the environment of some tryptophan residues changes on insertion and/or pore formation. In particular, spectroscopic anal. of a tryptophan mutant, W433F, suggests it is the residue mainly responsible for the obsd. effects. Furthermore, there is no change in the secondary structure content when the toxin inserts into membranes. Finally, the basis of the very low activity shown by a pneumolysin mol. from another strain of S. pneumoniae may be due to the movements of a key domain-domain interface. The mol. basis of pneumolysin -induced complement activation may be related to the structural similarity of one of the domains of pneumolysin to Fc, rather than the presumed homol. of the toxin to C-reactive protein as previously suggested. (c) 1998 Academic Press.

L3 ANSWER 4 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1998:122808 CAPLUS

DOCUMENT NUMBER:

128:240572

TITLE:

3

A conserved tryptophan in pneumolysin

is a determinant of the characteristics of

channels formed by pneumolysin in cells and planar lipid bilayers

AUTHOR(S):

Korchev, Yuri E.; Bashford, C. Lindsay; Pederzolli, Cecilia; Pasternak, Charles A.; Morgan, Peter J.; Andrew, Peter W.; Mitchell,

Timothy J.

CORPORATE SOURCE:

Department of Cellular and Molecular Sciences, St. George's Hospital Medical School, London,

SW17 ORE, UK

SOURCE:

Biochem. J. (1998), 329(3), 571-577

CODEN: BIJOAK; ISSN: 0264-6021

PUBLISHER:

Portland Press Ltd.

DOCUMENT TYPE:

Journal

LANGUAGE: English

Pneumolysin is one of the family of thiol-activatable, cytolytic toxins. Within these toxins the amino acid sequence Trp-Glu-Trp-Trp is conserved. Mutations made in this region of pneumolysin, residues 433-436 inclusive, did not affect cell binding or the formation of toxin oligomers in the target cell membrane. However, the mutations did affect hemolysis, leakage of low-mol.-mass metabolites from Lettre cells and the induction of conductance channels across planar lipid bilayers. Of eight modified pneumolysins examd., Trp-433 .fwdarw. Phe showed the smallest amt. of hemolysis or leakage (less than 5% of wild type). Pneumolysin -induced leakage from Lettre cells was sensitive to inhibition by bivalent cations but the extent of inhibition varied depending on the modification. Leakage by the mutant Trp-433 .fwdarw. Phe was least sensitive to cation inhibition. ion-conducting channels formed across planar lipid bilayers exhibit small (less than 30 pS), medium (30 pS-1 nS) and large (more than 1 nS) conductance steps. Small- and medium-sized channels were preferentially closed by bivalent cations. In contrast with wild-type toxin, which formed predominantly small channels, the modified toxin Trp-433 .fwdarw. Phe formed large channels that were insensitive to cation-induced closure. Polysaccharides of mol. mass more than 15 kDa inhibited hemolysis by wild-type toxin, but polysaccharide of up to 40 kDa did not prevent hemolysis by Trp-433 .fwdarw. Phe. Electron microscopy revealed that Trp-433 .fwdarw. Phe formed oligomeric arc and ring structures with dimensions identical with those of wild-type toxin, and that the ratio of arcs to rings formed was the same for wild-type toxin and the Trp-433 .fwdarw. Phe variant. We conclude that the change Trp-433 .fwdarw. Phe affects channel formation at a point subsequent to binding to the cell membrane and the formation of oligomers, and Searcher : Shears

that the size of arc and ring structures revealed by electron microscopy does not reflect the functional state of the channels.

L3ANSWER 5 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1996:201910 CAPLUS

DOCUMENT NUMBER:

124:255575

TITLE:

In vitro activities of U-100592 and U-100766,

novel oxazolidinone antibacterial agents

AUTHOR (S):

Zurenko, Gary E.; Yagi, Betty H.; Schaadt, Ronda

D.; Allison, John W.; Kilburn, James O.; Glickman, Suzanne E.; Hutchinson, Douglas K.; Barbachyn, Michael R.; Brickner, Steven J.

CORPORATE SOURCE:

Pharmacia & Upjohn, Inc., Kalamazoo, MI, 49001,

SOURCE:

Antimicrob. Agents Chemother. (1996), 40(4),

839-45

CODEN: AMACCQ; ISSN: 0066-4804

DOCUMENT TYPE:

Journal

LANGUAGE:

AΒ

English

Oxazolidinones make up a relatively new class of antimicrobial agents which possess a unique mechanism of bacterial protein synthesis inhibition. U-100592 $\{(S)-N-[[3-[3-fluoro-4-[4-in]]]\}$ (hydroxyacetyl) -1-piperazinyl]phenyl] -2-oxo-5-oxazolidinyl]methyl] acetamide and U-100766 { (S) -N-[[3-[3-fluoro-4-(4morpholinyl)phenyl]-2-oxo-5-oxazolidinyl]methyl]-acetamide} are novel oxazolidinone analogs from a directed chem. modification program. MICs were detd. for a variety of bacterial clin. isolates; the resp. MICs of U-100592 and U-100766 at which 90% of isolates are inhibited were as follows: methicillin-susceptible Staphylococcus aureus, 4 and 4 .mu.g/mL; methicillin-resistant S. aureus, 4 and 4 .mu.g/mL; methicillin-susceptible Staphylococcus epidermidis, 2 and 2 .mu.g/mL; methicillin-resistant S. epidermidis, 1 and 2 .mu.g/mL; Enterococcus faecalis, 2 and 4 .mu.g/mL; Enterococcus faecium, 2 and 4 .mu.g/mL; Streptococcus pyogenes, 1 and 2 .mu.g/mL; Streptococcus pneumoniae, 0.50 and 1 .mu.g/mL; Corynebacterium spp., 0.50 and 0.50 .mu.g/mL; Moraxella catarrhalis, 4 and 4 .mu.g/mL; Listeria monocytogenes, 8 and 2 .mu.g/mL; and Bacteroides fragilis, 16 and 4 .mu.g/mL. Most strains of Mycobacterium tuberculosis and the gram-pos. anaerobes were inhibited in the range of 0.50 to 2 .mu.g/mL. Enterococcal strains resistant to vancomycin (VanA, VanB, and VanC resistance phenotypes), pneumococcal strains resistant to penicillin, and M. tuberculosis strains resistant to common antitubercular agents (isoniazid, streptomycin, rifampin, ethionamide, and ethambutol) were not cross-resistant to the oxazolidinones. The presence of 10, 20, and 40% pooled human serum did not affect the antibacterial activities of the oxazolidinones. Time-kill studies demonstrated a bacteriostatic effect of the analogs against staphylococci and enterococci but a bactericidal Searcher : Shears 308-4994

effect against streptococci. The spontaneous mutation frequencies of S. aureus ATCC 29213 were <3.8 .times. 10-10 and <8 .times. 10-11 for U-100592 and U-100766, resp. Serial transfer of three staphylococcal and two enterococcal strains on drug gradient plates produced no evidence of rapid resistance development. Thus, these new oxazolidinone analogs demonstrated in vitro antibacterial activities against a variety of clin. important human pathogens.

ANSWER 6 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1993:488721 CAPLUS

DOCUMENT NUMBER:

119:88721

TITLE:

Ethanol impairs neutrophil chemotaxis in vitro but not adherence or recruitment to lungs of

rats with experimental pneumococcal

pneumonia

AUTHOR (S):

Lister, Philip D.; Gentry, Martha J.; Preheim,

Laurel C.

CORPORATE SOURCE:

Sect. Infect. Dis., VA Med. Cent., Omaha, NE,

USA

SOURCE:

J. Infect. Dis. (1993), 167(5), 1131-7

CODEN: JIDIAQ; ISSN: 0022-1899

DOCUMENT TYPE:

Journal

LANGUAGE:

English

AB The effect of 7 days of ethanol ingestion on circulating neutrophil (PMNL) counts and PMNL adherence, chemotaxis, and recruitment was investigated. Pair-feeding of rats resulted in a significant decrease in PMNL counts in both ethanol-fed and control rats. mean no. of PMNL exhibiting chemotaxis in a modified Boyden chamber in response to lipopolysaccharide-activated normal rat serum was significantly decreased in ethanol-fed rats compared with controls. The percentage of adherence to nylon wool columns, however, was similar in both groups. To measure pulmonary PMNL recruitment, rats were infected transtracheally with 105 Streptococcus pneumoniae and sacrificed. Bronchoalveolar lavage fluid from both groups contained similar nos. of PMNL 8 h after infection. By 24 h, PMNL nos. in lavage fluid from ethanol-fed rats exceeded those in controls. PMNL recruitment continued in the ethanol-fed rats at 48 and 72 h, whereas values in controls had returned to baseline. Thus, the impaired pulmonary defense against S. pneumoniae in ethanol-fed rats is not due to defective PMNL recruitment.

ANSWER 7 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1992:509828 CAPLUS

DOCUMENT NUMBER:

117:109828

TITLE:

Comparative efficacy of autolysin and pneumolysin as immunogens protecting mice against infection by Streptococcus

pneumoniae

Searcher :

Shears 308-4994

AUTHOR (S):

Lock, Robert A.; Hansman, David; Paton, James C.

CORPORATE SOURCE:

Dep. Microbiol., Adelaide Child. Hosp., North

Adelaide, Australia

SOURCE:

Microb. Pathog. (1992), 12(2), 137-43

CODEN: MIPAEV; ISSN: 0882-4010

DOCUMENT TYPE:

Journal English

LANGUAGE: AB

Previous studies on S. pneumoniae have established that the pneumococcal proteins autolysin (N-acetylmuramyl-L-alanine amidase) and pneumolysin both contribute significantly to the virulence of the organism. Here, autolysin and a defined toxoid deriv. of pneumolysin were tested, individually and in combination, for efficacy in a mouse model as antigens protecting against challenge with virulent, wild-type S. pneumoniae. While each antigen alone provided significant protection, the degree of protection was not increased when the antigens were administered together. In an addnl. expt., mice were challenged with a genetically-modified mutant strain of pneumococcus unable to express active pneumolysin.

Pre-immunization of such mice with autolysin failed to provide any significant protection against the challenge. Apparently, the most important contribution made by autolysin to the virulence of S. pneumoniae may be its role in mediating the release of pneumolysin from the pneumococcal cytoplasm during infection.

ANSWER 8 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1992:122142 CAPLUS

DOCUMENT NUMBER:

116:122142

TITLE:

Characterization and sequence of the Escherichia

coli stress-induced psp operon

AUTHOR (S):

Brissette, Janice L.; Weiner, Lorin; Ripmaster,

Tracy L.; Model, Peter

CORPORATE SOURCE:

Rockefeller Univ., New York, NY, 10021, USA

J. Mol. Biol. (1991), 220(1), 35-48

CODEN: JMOBAK; ISSN: 0022-2836

DOCUMENT TYPE:

Journal

LANGUAGE:

SOURCE:

English

AB A new E. coli operon, the phage shock protein (psp) operon, is induced in response to heat, ethanol, osmotic shock and infection by filamentous bacteriophages. The operon includes at least four genes: pspA, B, C and E. PspA assocs. with the inner membrane and has the heptad repeats characteristic of proteins that can form coiled coils. The operon encodes a factor that activates psp expression, and deletion analyses indicate that this protein is PspC; PspC is predicted to possess a leucine zipper, a motif present in many eukaryotic transcription factors. The pspE gene is expressed in response to stress as part of the operon, but is also transcribed from its own promoter under normal conditions. In vitro

Searcher : Shears

studies suggest that PspA and C are modified in vivo. Expression of the psp genes does not require the heat shock sigma factor, .sigma.32. The increased duration of psp induction in a .sigma.32 mutant suggests that a product (or products) of the heat shock response down-regulates expression of the operon.

1.3 ANSWER 9 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1992:103557 CAPLUS

DOCUMENT NUMBER:

116:103557

TITLE:

Pneumolysin induces the salient histologic features of pneumococcal infection in the rat lung in vivo

AUTHOR (S):

Feldman, C.; Munro, N. C.; Jeffery, P. K.;

Mitchell, T. J.; Andrew, P. W.; Boulnois, G. J.; Guerreiro, D.; Rohde, J. A. L.; Todd, H. C.; et

al.

CORPORATE SOURCE:

Dep. Thorac. Med., Natl. Heart Lung Inst.,

London, SW3 6LR, UK

SOURCE:

Am. J. Respir. Cell Mol. Biol. (1991), 5(5),

416-23

CODEN: AJRBEL; ISSN: 1044-1549

DOCUMENT TYPE:

Journal

LANGUAGE:

English AB Streptococcus pneumoniae infections are common, but how they cause host tissue injury and death is incompletely understood. Immunization with pneumolysin, a thiol-activated toxin produced by the pneumococcus, partially protects animals during subsequent infection. The mechanism by which pneumolysin contributes to disease is not known. The aim of the present investigation was to det. the hitol. changes induced by recombinant pneumolysin in the rat lung and to compare them with the changes induced by live organisms. Injection of either toxin (200 or 800 ng) or bacteria into the apical lobe bronchus was assocd. with the development of a severe lobar pneumonia restricted to the apical lobe. The changes induced by the toxin were greater at the higher concn., and changes were most severe in those animals in which there was partial ligation of the apical lobe bronchus. The pneumonitis was less severe following injection of a modified toxin with decreased hemolytic activity, generated by site-directed mutagenesis of the cloned pneumolysin gene, indicating that this property of the toxin was important in generating pulmonary inflammation. was still considerable pneumonitis after injection of a

ANSWER 10 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

complement.

1992:77475 CAPLUS

DOCUMENT NUMBER:

116:77475

modified toxin with decreased capacity to activate

Searcher :

Shears 308-4994 TITLE: Truncated forms of PspA that are secreted from

Streptococcus pneumoniae and their use in

functional studies and cloning of the pspA gene

AUTHOR (S): Yother, Janet; Handsome, Geraline L.; Briles,

David E.

CORPORATE SOURCE: Dep. Microbiol., Univ. Alabama, Birmingham, AL,

35294, USA

SOURCE: J. Bacteriol. (1992), 174(2), 610-18

CODEN: JOBAAY; ISSN: 0021-9193

DOCUMENT TYPE:

Journal

LANGUAGE: English

Insertion-duplication mutagenesis was used to generate mutants of S. pneumoniae that produced truncated forms of PspA (pneumococcal surface protein A). The truncated products, representing from 20 to 80% of the complete PspA mol., were all secreted from the cell and could be detected in unconcd. culture medium. Anal. of the truncated mols. showed that the antigenic variability known to be assocd. with PspA is located in the .alpha.-helical N-terminal half of the mol. This region was also found to contain immunogenic and protection-eliciting epitopes and to define the max. region of the mol. that is likely to be surface exposed. The apparent mol. wt. variability seen for PspA mols. of different S. pneumoniae strains was localized to both the N- and C-terminal halves of the protein. Attachment of PspA to S. pneumoniae was found to require regions located carboxy to the 5th repeat unit in the C-terminal end of the mol. From the insertion-duplication mutants, the complete pspA gene was cloned and expressed in Escherichia coli. Differences in apparent mol. wt. were obsd. when the same cloned product was expressed in E. coli and S. pneumoniae, suggesting that PspA is modified differently in the 2 hosts.

ANSWER 11 OF 26 CAPLUS COPYRIGHT 1999 ACS L3

ACCESSION NUMBER: 1991:654020 CAPLUS

DOCUMENT NUMBER: 115:254020

TITLE: Complement activation and antibody binding by

pneumolysin via a region of the toxin

homologous to a human acute-phase protein

AUTHOR (S): Mitchell, T. J.; Andrew, P. W.; Saunders, F. K.;

Smith, A. N.; Boulnois, G. J.

CORPORATE SOURCE: Dep. Microbiol., Univ. Leicester, Leicester, LE1

9NH, UK

SOURCE: Mol. Microbiol. (1991), 5(8), 1883-8

CODEN: MOMIEE; ISSN: 0950-382X

DOCUMENT TYPE:

Journal

LANGUAGE: English

Pneumolysin, a membrane-damaging toxin, is known to activate the classical complement pathway. It was shown that 1 .mu.g mL-1 of penumolysin can activate complement, which is a much

Searcher Shears :

lower level than obsd. previously. Two distinct regions of pneumolysin were identified which show homol. with a contiguous sequence within acute-phase proteins, including human C-reactive protein (CRP). Site-directed mutagenesis of the pneumolysin gene was used to change residues common to pneumolysin and CRP. Some of the modified toxins had a reduced ability both to activate complement and bind antibody. It is suggested that the ability of pneumolysin to activate complement is related to its ability to bind the Fc portion of IqG.

ANSWER 12 OF 26 CAPLUS COPYRIGHT 1999 ACS L3

ACCESSION NUMBER:

1989:528664 CAPLUS

DOCUMENT NUMBER:

111:128664

TITLE:

Pneumolysin, the thiol-activated toxin

of Streptococcus pneumoniae, does not require a

thiol group for in vitro activity

AUTHOR (S):

Saunders, F. K.; Mitchell, T. J.; Walker, J. A.;

Andrew, P. W.; Boulnois, G. J.

CORPORATE SOURCE:

Dep. Microbiol., Univ. Leicester, Leicester, LE1

9HN, UK

SOURCE:

Infect. Immun. (1989), 57(8), 2547-52

CODEN: INFIBR; ISSN: 0019-9567

DOCUMENT TYPE:

Journal

LANGUAGE: English

The role of the single cysteine residue in the activity of the ΔR thiol-activated toxin pneumolysin was investigated using oligonucleotide-mediated, site-directed mutagenesis. Three modified toxins in which the cysteine residue was changed in an alanine, a serine, or a glycine residue were purified to homogeneity and examd. for activity. The Cys-428.fwdarw.Ala modified toxin was indistinguishable from the wild-type recombinant toxin in terms of hemolytic activity and lytic and inhibitory effects on human polymorphonuclear leukocytes (PMN), indicating that the cysteine residue is not essential for toxin activity. The Cys-428.fwdarw.Ser and Cys-429.fwdarw.Gly modified toxins had reduced activity on erythrocytes and polymorphonuclear leukocytes, being 6 and 20 times less active than the wild type, resp. However, all the modified toxins formed oligomers in erythrocyte membranes to the same extent as the wild-type recombinant toxin. This suggests that the cysteine residue at position 428 is involved in neither the binding of toxin to membranes nor its insertion into the membrane, and also that the formation of oligomers is not by itself sufficient for toxin activity.

ANSWER 13 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1988:547401 CAPLUS

DOCUMENT NUMBER:

109:147401

Searcher :

Shears 308-4994 TITLE:

Stimulation of human neutrophils, monocytes, and

platelets by modified C-reactive

protein (CRP) expressing a neoantigenic

specificity

AUTHOR (S):

Potempa, L. A.; Zeller, J. M.; Fiedel, B. A.;

Kinoshita, C. M.; Gewurz, H.

CORPORATE SOURCE:

Dep. Immunol./Microbiol., Rush Presbyterian St.

Luke's Med. Cent., Chicago, IL, USA

SOURCE:

Inflammation (N. Y.) (1988), 12(4), 391-405

CODEN: INFLD4; ISSN: 0360-3997

DOCUMENT TYPE:

LANGUAGE:

Journal English

C-reactive protein (CRP) can be structurally modified by heat, acid, or urea-chelation to express a neoantigen designated as neo-CRP. This antigen is also expressed on the in vitro primary protein translocation products of both human and rabbit CRP. Unmodified CRP and CRP complexed with pneumococcal C-polysaccharide (CPS) do not express neo-CRP. Forms of CRP expressing neo-CRP but not native CRP antigenicity (even in the presence of CPS) consistently and in a dose-dependent manner potentiated the respiratory burst response of human polymorphonuclear leukocytes and peripheral blood monocytes to heat-modified IgG. Forms of CRP expressing neo-CRP antigenicity also induced reactions of aggregation and secretion from isolated platelets and potentiated platelet activation stimulated by ADP in platelet-rich-plasma, while native CRP alone or complexed with CPS again did not. Unlike CRP-CPS complexes, forms of CRP expressing neo-CRP were not able to activate the complement system. These data emphasize the biol. potential inherent in this humoral acute-phase reactant, particularly in the activation of the formed elements of the blood important in the inflammatory response. Since these cell-activating properties are preferentially obsd. when CRP is structurally modified to express the neo-CRP antigen, such a mol. conversion may be central to the structure-function relationships of CRP at local sites of inflammation and tissue injury.

L3 ANSWER 14 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1988:17214 CAPLUS

DOCUMENT NUMBER:

108:17214

TITLE:

Overproduction and rapid purification of the

amidase of Streptococcus pneumoniae

AUTHOR(S):

Garcia, J. L.; Garcia, E.; Lopez, R.

CORPORATE SOURCE:

Cent. Invest. Biol., C.S.I.C., Madrid, 28006,

Spain

SOURCE:

Arch. Microbiol. (1987), 149(1), 52-6

CODEN: AMICCW; ISSN: 0302-8933

DOCUMENT TYPE:

Journal

LANGUAGE:

English Searcher: Shears 308-4994 AB Oligonucleotide-directed mutagenesis of a plasmid contg. the lytA gene coding for the pneumococcal amidase has allowed the sepn. of the coding sequence of the gene. This sequence has been placed in plasmid pIN-III(lppP-5)-A3 downstream from both a modified lipoprotein promoter and the lactose promoter to construct the recombinant plasmid pGL100. When Escherichia coli RB 791 (pGL100) was grown in the presence of lactose, the pneumococcal amidase accounted for 7% of the total protein present in this strain after 18 h incubation at 37.degree.. overproduced amidase was purified in a single-step procedure using a choline-Sepharose 6B column, taking advantage of the fact that this enzyme was the unique protein with affinity for choline present in exts. obtained from E. coli RB791 (pGL100). The development of the above design opens up the possibility of studying the mechanism that regulates the activity of this important autolysin by using physicochem. techniques that require the availability of high amts. of purified amidase.

L3 ANSWER 15 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1985:519684 CAPLUS

DOCUMENT NUMBER:

103:119684

TITLE:

Localized conversion in Streptococcus pneumoniae

recombination: heteroduplex preference

AUTHOR (S):

Sicard, Michel; Lefevre, Jean Claude; Mostachfi,

Pezechpour; Gasc, Anne Marie; Sarda, Claudine

CORPORATE SOURCE:

Cent. Rech. Biochim. Genet. Cell., CNRS,

Toulouse, 31062, Fr.

SOURCE:

Genetics (1985), 110(4), 557-68 CODEN: GENTAE; ISSN: 0016-6731

DOCUMENT TYPE:

Journal English

LANGUAGE:

In pneumococcal transformation, the frequency of recombinants between point mutations is generally proportional to distance. An aberrant marker in the amiA locus enhanced recombination frequency when crossed with any other allele of this gene. The hyperrecombination obsd. in 2-point crosses could be explained by 2 hypotheses: the aberrant marker induces frequent crossovers in its vicinity or the mutant is converted to wild type. Evidence is presented to show that in suitable 3-point crosses, this hyperrecombination does not modify the recombination frequency between outside markers, suggesting that a conversion occurs at the site of this mutation. To est. the length over which this event occurs, very closely linked markers were isolated and used in 2-point crosses. It appears that the conversion system removes only a few base pairs (from 3-27) around the aberrant marker. This conversion process is quite different from the mismatch-repair system controlled by hex genes in pneumococcus, which involves several thousand base pairs. Moreover, artificial heteroduplexes were constructed using sepd. DNA

strands. It appears that only 1 of the 2 heteroduplexes is specifically converted. The conversion system acts upon 5'..ATTAAT..3'/3'..TAAGTA..5'. A possible role of the palindrome resulting from the mutation is discussed.

ANSWER 16 OF 26 CAPLUS COPYRIGHT 1999 ACS L3

ACCESSION NUMBER:

1985:22561 CAPLUS

DOCUMENT NUMBER:

102:22561

TITLE:

Opsonic activity of immunoglobulin prepared for

intravenous use

AUTHOR (S):

Hetherington, Seth V.; Giebink, G. Scott

CORPORATE SOURCE:

Dep. Pediatr., Albany Med. Cent., Albany, NY,

12208, USA

SOURCE:

J. Lab. Clin. Med. (1984), 104(6), 977-86

CODEN: JLCMAK; ISSN: 0022-2143

DOCUMENT TYPE:

Journal English

LANGUAGE:

AΒ

The opsonic activity of 2 Ig prepns. modified for i.v.

infusion was tested against Streptococcus pneumoniae types 3, 7F,

and 14 and 2 strains of Staphylococcus aureus by polymorphonuclear leukocyte uptake of [3H]thymidine-labeled bacteria. Reduced and alkylated Ig (Chem-IgG) and Ig prepd. by chromatog. with diethylaminoethyl-Sephadex (DEAE-IgG) were evaluated with and without complement and compared with the opsonic activity of immune serum globulin and heated pooled human serum. Opsonic activity of DEAE-IgG was greater than that of Chem-IgG and equiv. to the activity of immune serum globulin and pooled human serum against S. aureus 502A and type 3 pneumococcus. Both i.v. Igs had lower opsonic activity than either pooled human serum or immune serum globulin against type 14 pneumococcus. There were no differences in antibody avidity for pneumococcal antigen among the Igs tested. All 4 opsonins had similar opsonic activity against the protein A-deficient S. aureus Wood 46. Modification of Ig for i.v. infusion by chem. alteration may

adversely affect opsonic activity by changing the Fc portion of the antibody mol.

ANSWER 17 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1984:101355 CAPLUS

DOCUMENT NUMBER:

100:101355

TITLE:

The role of C5 in polymorphonuclear leukocyte recruitment in response to

Streptococcus pneumoniae

AUTHOR (S):

Toews. Galen B.; Vial, Wayne C.

CORPORATE SOURCE:

Southwestern Med. Sch., Univ. Texas, Dallas, TX,

75235, USA

Journal

SOURCE:

Am. Rev. Respir. Dis. (1984), 129(1), 82-6

CODEN: ARDSBL; ISSN: 0003-0805

DOCUMENT TYPE:

LANGUAGE:

English

AB

Congenic complement C5-sufficient B10.D2/nSn (C5+) and C5-deficient B10.D2/oSn (C5-) mice were used to det. the importance of the C5 mol. in the polymorphonuclear leukocyte (PMN) response to S. pneumoniae. The C5+ and C5- mice were injected with water and varying inoculums of pneumococci via an endobronchial catheter. Bronchoalveolar lavage (BAL) was performed on the inoculated lung at 0 and 4 h after injection. Cellular response was measured and chemotactic activity was assayed in BAL supernatants at each time interval, using human PMN in a modified Boyden chamber. Clearance of bacteria was studied by a quant. lung The C5+ mice recruited more PMN after challenges with both culture. 104 and 106 pneumococci than did the C5- mice; however, PMN accumulation did occur in C5- mice. Similarly, C5+ mice generated more intraalveolar chemotactic activity than did C5- mice, but chemotactic activity was present in both C5+ and C5- mice, as detd. by checkerboard assays. Pulmonary clearance of bacteria was impaired in the absence of C5 at both inoculums. Thus, the C5 mol. yields important PMN chemotaxins during the early time period after intrapulmonary inoculation of S. pneumoniae. However, PMN recruitment after this insult also results from other chemotaxins since both chemotactic activity and PMN recruitment occur within the alveoli of C5- mice.

ANSWER 18 OF 26 CAPLUS COPYRIGHT 1999 ACS L3

ACCESSION NUMBER:

1983:102390 CAPLUS

DOCUMENT NUMBER:

98:102390

TITLE:

Mutagenesis in Streptococcus pneumoniae (Pneumococcus) by

transformation with DNA modified by

the carcinogen-mutagen, aflatoxin B1 Stark, Avishay A.; Giroux, Craig N.

AUTHOR (S):

Dep. Biochem., Tel-Aviv Univ., Tel-Aviv, Israel

CORPORATE SOURCE: SOURCE:

Mutat. Res. (1983), 107(1), 23-32

CODEN: MUREAV; ISSN: 0027-5107

DOCUMENT TYPE:

Journal

LANGUAGE:

English

GΙ

Searcher

Ι

Shears 308-4994 AB Covalent binding of metabolically activated aflatoxin B1 (AFB1)(I) [1162-65-8] to transforming DNA of Pneumococcus caused a decrease in transforming activity and induced mutations to antibiotic resistance in recipients. The 2 effects were proportional, in vitro, to the covalent binding levels of AFB1 to pneumococcal DNA.

L3 ANSWER 19 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1975:560665 CAPLUS

DOCUMENT NUMBER:

83:160665

TITLE:

Modifier mutation affecting

AUTHOR (S): CORPORATE SOURCE: utilization of mannitol in pneumococcus Rotheim, Minna B.; Hotchkiss, Rollin D.

Upstate Med. Cent., State Univ. New York,

Syracuse, N. Y., USA

SOURCE:

Can. J. Microbiol. (1975), 21(8), 1139-43

CODEN: CJMIAZ

DOCUMENT TYPE:

Journal

LANGUAGE:

English

AB The altered growth of a pneumococcal mutant contg. marker M, which confers ability to utilize mannitol, and a modifier gene is reported. The modifier gene is closely linked to the M gene and imposes a requirement for growth in 0.4% glucose before growth in mannitol medium. The tentative position of the M gene of pneumococcus is ery-r str-r M sul-rd.

L3 ANSWER 20 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1974:548335 CAPLUS

DOCUMENT NUMBER:

81:148335

TITLE:

Marker discrimination and mutagen

induced alterations in pneumococcal

transformation

AUTHOR (S):

Tiraby, Jean G.; Fox, Maurice S.

CORPORATE SOURCE:

Dep. Biol., Massachusetts Inst. Technol.,

Cambridge, Mass., USA

SOURCE:

Genetics (1974), 77(3), 449-58

CODEN: GENTAE

DOCUMENT TYPE:

Journal

LANGUAGE:

English

Nitrous acid-induced and NH2OH-induced chem. alterations in transforming DNA resulted in the loss of biol. activity and in mutagenesis. The function responsible for the discrimination between high efficiency and low efficiency markers in pneumococcal transformation, and for the elimination of a substantial fraction of spontaneously occurring mutational events, did not appear to act on integrated DNA carrying these chem.

alterations. The chem. modifications resulted in mutations that were evident among bacteria transformed with the treated DNA. Fusidic acid-resistant mutants isolated in this way were predominantly of the low efficiency class. Most mutations of spontaneous origin occurring in this locus are of the high efficiency class. Discrimination probably results from the elimination of specific classes of base pair mismatches that occur as intermediates in transformation. The base pair mismatches most effectively eliminated in the discriminating strain of pneumococcus are probably those of the A:C and G:T type and the immediate product of transformation with mutagenized DNA involves intermediates that are not recognized as A:C or G:T by the discrimination system.

L3 ANSWER 21 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER: 1974:35024 CAPLUS

DOCUMENT NUMBER: 80:35024

TITLE: Integration efficiency in DNA-induced

transformation of Pneumococcus

[Diplococcus pneumoniae]. II. Genetic studies

of mutant integrating all the markers

with a high efficiency

AUTHOR(S): Tiraby, Gerard; Sicard, Michel A.

CORPORATE SOURCE: Lab. Genet., Univ. Paul Sabatier, Toulouse, Fr.

SOURCE: Genetics (1973), 75(1), 35-48

CODEN: GENTAE

DOCUMENT TYPE: Journal LANGUAGE: English

AB Transformation of the D. pneumosiae mutant 401 by DNA's bearing a std. ref. marker and several other markers belonging to 2 unlinked loci showed that differences in the integration efficiencies of these markers were considerably reduced in this strain compared to the wild-type strain Cl3. The sensitivities of mutant 401 to uv light and to x-irradn. were the same as those of Cl3. However, in 401 all the markers tested were more resistant to inactivation as shown by transformation of 401 and Cl3 by uv-irradiated DNA. The increase in resistance was greater for low-efficiency (LE) markers than for high-efficiency (HE) markers. The decreased discrimination between LE and HE markers in strain 401 was not due to a mechanism related to modification of markers in the transforming DNA by the recipient cells, nor were the proteins inducing competence of the cells responsible for the differences in the integration efficiencies of various markers. Genetic studies of the fate of recombinants as well as the measure of the amt. of DNA taken up showed that all markers were integrated into strain 401 by the same recombination process, that specific for HE markers.

ACCESSION NUMBER:

1974:35023 CAPLUS

DOCUMENT NUMBER:

80:35023

TITLE:

Integration efficiency in DNA-induced

transformation of pneumococcus

[Diplococcus pneumoniae]. I. Method of

transformation in solid medium and its use for

isolation of transformation-deficient and

recombination-modified mutants

AUTHOR (S):

Tiraby, Gerard; Claverys, Jean P.; Sicard,

Michel A.

CORPORATE SOURCE:

Lab. Genet., Univ. Paul Sabatier, Toulouse, Fr.

SOURCE:

Genetics (1973), 75(1), 23-33

CODEN: GENTAE

DOCUMENT TYPE:

Journal

LANGUAGE:

English

A method of transformation on solid medium esp. adapted for D. AΒ pneumoniae was developed. Under specific conditions, all colonies that were allowed to grow in the presence of transforming DNA gave rise to transformed bacteria. Combined with replica plating this technique was used to isolate mutants modified with regard to recombination. Most of the mutants found were transformation-defective and showed a large diversity in their response to uv light. Some of these mutants had lost their ability to take up transforming DNA. One mutant showed a reduced yield of transformants for a given quantity of DNA taken up. Mutants that manifest altered behavior with regard to marker efficiencies were also isolated. One of these exhibited a decrease in the transformation efficiency of only the high-efficiency markers and 2 mutants showed a decrease in the transformation efficiency of the low efficiency markers.

L3 ANSWER 23 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1970:9696 CAPLUS

DOCUMENT NUMBER:

72:9696

TITLE:

Comparison of the x-ray sensitivities of

thymidine, transforming DNA, and bacteria

AUTHOR (S):

Apelgot, Sonia; Rebeyrotte, N.

CORPORATE SOURCE:

Lab. Curie Pasteur, Inst. Radium, Paris, Fr.

SOURCE:

Biophysik (1969), 6(1), 25-33

CODEN: BPYKAU

DOCUMENT TYPE:

Journal

LANGUAGE:

French

AB Thymidine-6-3H (I), transforming DNA (II) isolated from a culture of pneumococcus R36A Smr (a mutant resistant to streptomycin), Escherichia coli strain B3 thy- Smr with high radioresistance, and Diplococcus pneumoniae strain R36A Smr were irradiated. The bacteria were not very sensitive to irradn., but I and II were. The sensitivity varied with the temp. and media used during the irradn. A graph of the ratio of I decomposed when Searcher: Shears 308-4994

irradiated to the value for I irradiated vs. the irradn. temp. was an upward concave curve. A graph of the ratio of the value of x-ray doses which left a 37% survival of bacteria or II irradiated at 0.degree. in saline to the dose needed with oth er exptl. conditions of temp. and media vs. the irradn. temp. was a downward concave curve. The weak dependence of bacterial radiosensitivity to environmental modifications is a property of cells. The effect of lowering temp. seemed to be a property of the DNA, not the bacteria.

L3 ANSWER 24 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1969:458006 CAPLUS

DOCUMENT NUMBER:

71:58006

TITLE:

Genetic analysis by transformation of a group of

uracil requiring mutants of Diplocococcus pneumoniae

AUTHOR (S):

Morse, Helvise G. G.; Lerman, L. S.

CORPORATE SOURCE:

Med. Center, Univ. of Colorado, Denver, Colo.,

USA

SOURCE:

Genetics (1969), 61(1), 41-60

CODEN: GENTAE

DOCUMENT TYPE:

Journal English

LANGUAGE:

A no. of auxotrophic pneumococcus mutants were

isolated from a modified medium. Mutants
requiring uracil, lysine, and phenylalanine were recovered. Twelve
of the 40 uracil-requiring mutants were used in
recombination expts. The transformation yield in genetic crosses
was dependent on the efficiency of transformation of the recipient
mutant. When allowance was made for recipient variation in
a recombination index, reciprocals were closely related to each
other, and a linear arrangement of linked markers was possible. The
comparison of long map distances with the sum of intervening short
distances indicated that neg. interference was operative in regions
<0.2 index unit. Reasonably good additivity among recombination
indexes occurred, and was improved by correction for neg.

L3 ANSWER 25 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

interference.

1968:112427 CAPLUS

DOCUMENT NUMBER:

68:112427

TITLE:

Role of integrity and heterogeneity of

deoxyribonucleic acid molecules during genetic

recombination in pneumococcal

transformation

AUTHOR(S):

Kent, Joan L.; Hotchkiss, Rollin D.

CORPORATE SOURCE:

Dep. Genet., Rockefeller Inst., New York, N. Y.,

USA

SOURCE:

Physiol. Gene Mutat. Expression, Proc. Symp.

(1966), Meeting Date 1965, 165-71

CODEN: 19PXAO

DOCUMENT TYPE:

Conference

LANGUAGE:

AΒ

English

The effect of phys. and chem. modified DNA mols. on genetic recombination was investigated in various pneumococcal mutants. A method was developed which allowed the detn. of the amt. of transformation caused by the exposure to DNA for various time intervals. For mutants with 1 marker (e.g., streptomycin resistance or sulfonamide resistance) a linear relation was found between the time of exposure and the amt. of transformation. The rate for any given marker was detd. by the DNA concn. Two markers were often incorporated together or linked. The 3-factor linkage group was exploited by studying the effect of chem. modification of the DNA on single, multiple, and complex transformant classes. The kinetic study on linkage of DNA denatured by heating and renatured by cooling showed that the helical structure of DNA was of importance in the penetration step. Denaturated DNA competed with native DNA. In further expts. the technique of subcrit. heat inactivation of a multiply marked DNA was used. The decay of transformations (assayed on a concd., wild-type culture at 3 .times. 107 viable counts/ml. exposed to 0.02 .mu.g. of modified DNA for 5 min. at 30.degree.) accumulated more quickly in the case of a triply marked group than in a singly marked one. Tightly linked markers, loosely linked markers, and unlike pairs were usually inactivated at the same exponential rate for a typical single marker. More interestingly, the cells seemed to be able to escape the consequences of exposure to damaged DNA. If the damaged sites were incorporated one would expect either a mutagenic or a lethal effect. Neither one was observed in the appropriate expts.

L3 ANSWER 26 OF 26 CAPLUS COPYRIGHT 1999 ACS

ACCESSION NUMBER:

1967:616 CAPLUS

DOCUMENT NUMBER:

66:616

TITLE:

Interaction of mutations affecting

growth rate and resistance to streptomycin in

pneumococci and streptococci

AUTHOR (S):

SOURCE:

Krauss, Marjorie R.; King, James Clement; Cox,

Rody P.

CORPORATE SOURCE:

New York Univ. Med Center, New York, N. Y., USA

J. Bacteriol. (1966), 92(5), 1337-44

CODEN: JOBAAY

DOCUMENT TYPE:

Journal

LANGUAGE:

English

AB A strain of Streptococcus (viridans group) was shown by transformation reactions to be the carrier of 2 interacting mutations. One produced resistance to streptomycin and a slow rate of growth; the only effect of the 2nd was an increase in

growth rate when it was added by transformation to streptococcal strains that had already been transformed to bear the 1st. Similar modifying mutations were observed in strains of streptococci and pneumococci into which the 1st mutation had been introduced by transformation. 16 references.

=> d his 14-; d 1-27 ibib abs

(FILE 'MEDLINE, BIOSIS, EMBASE, JICST-EPLUS, TOXLINE, TOXLIT, CONFSCI, PHIC, PHIN' ENTERED AT 16:04:40 ON 02 SEP 1999)

L4 80 S L3

L5 27 DUP REM L4 (53 DUPLICATES REMOVED)

L5 ANSWER 1 OF 27 MEDLINE DUPLICATE 1

ACCESSION NUMBER: 1998109732 MEDLINE

DOCUMENT NUMBER: 98109732

TITLE: A conserved tryptophan in pneumolysin is a

determinant of the characteristics of channels formed

by pneumolysin in cells and planar lipid

bilayers.

AUTHOR: Korchev Y E; Bashford C L; Pederzolli C; Pasternak C

A; Morgan P J; Andrew P W; Mitchell T J

CORPORATE SOURCE: Department of Cellular and Molecular Sciences, St.

George's Hospital Medical School, London, U.K.

SOURCE: BIOCHEMICAL JOURNAL, (1998 Feb 1) 329 (Pt 3) 571-7.

Journal code: 9YO. ISSN: 0264-6021.

PUB. COUNTRY: ENGLAND: United Kingdom

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals; Cancer Journals

ENTRY MONTH: 199805 ENTRY WEEK: 19980501

Pneumolysin is one of the family of thiol-activatable, AB cytolytic toxins. Within these toxins the amino acid sequence Trp-Glu-Trp-Trp is conserved. Mutations made in this region of pneumolysin, residues 433-436 inclusive, did not affect cell binding or the formation of toxin oligomers in the target cell membrane. However, the mutations did affect haemolysis, leakage of low-molecular-mass metabolites from Lettre cells and the induction of conductance channels across planar lipid bilayers. Of eight modified pneumolysins examined, Trp-433-->Phe showed the smallest amount of haemolysis or leakage (less than 5% of wild type). Pneumolysin-induced leakage from Lettre cells was sensitive to inhibition by bivalent cations but the extent of inhibition varied depending on the modification. Leakage by the mutant Trp-433-->Phe was least sensitive to cation inhibition. The ion-conducting

channels formed across planar lipid bilayers exhibit small (less

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than 30 pS), medium (30 pS-1 nS) and large (more than 1 nS) conductance steps. Small- and medium-sized channels were preferentially closed by bivalent cations. In contrast with wild-type toxin, which formed predominantly small channels, the modified toxin Trp-433-->Phe formed large channels that were insensitive to cation-induced closure. Polysaccharides of molecular mass more than 15 kDa inhibited haemolysis by wild-type toxin, but polysaccharide of up to 40 kDa did not prevent haemolysis by Trp-433-->Phe. Electron microscopy revealed that Trp-433-->Phe formed oligomeric arc and ring structures with dimensions identical with those of wild-type toxin, and that the ratio of arcs to rings formed was the same for wild-type toxin and the Trp-433-->Phe variant. We conclude that the change Trp-433-->Phe affects channel formation at a point subsequent to binding to the cell membrane and the formation of oligomers, and that the size of arc and ring structures revealed by electron microscopy does not reflect the functional state of the channels.

L5 ANSWER 2 OF 27 MEDLINE

DUPLICATE 2

ACCESSION NUMBER:

1999033058 99033058

DOCUMENT NUMBER:

The molecular mechanism of pneumolysin, a

MEDLINE

virulence factor from Streptococcus pneumoniae.

AUTHOR:

Rossjohn J; Gilbert R J; Crane D; Morgan P J; Mitchell T J; Rowe A J; Andrew P W; Paton J C; Tweten

R K; Parker M W

CORPORATE SOURCE:

The Ian Potter Foundation Protein Crystallography
Laboratory, St. Vincent's Institute of Medical
Research, 41 Victoria Parade, Fitzroy, Victoria,
3065, Australia.. jamie@brains.medstv.unimelb.edu.au
JOURNAL OF MOLECULAR BIOLOGY, (1998 Nov 27) 284 (2)

SOURCE:

449-61. Journal code: J6V. ISSN: 0022-2836.

PUB. COUNTRY:

ENGLAND: United Kingdom

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Priority Journals; Cancer Journals

ENTRY MONTH:

199902

ENTRY WEEK:

19990204

AB Pneumolysin, a member of the thiol-activated cytolysin family of toxins, is a virulence factor from the Gram-positive bacterium Streptococcus pneumoniae. The toxin forms large oligomeric pores in cholesterol-containing membranes of eukaryotic cells. A plethora of biochemical and mutagenesis data have been published on pneumolysin, since its initial characterization in the 1930s. Here we present an homology model of the monomeric and oligomeric forms of pneumolysin based on the recently determined crystal structure of perfringolysin O and electron microscopy data. A feature of the model is a striking

electronegative surface on parts of pneumolysin that may reflect its cytosolic location in the bacterial cell. The models provide a molecular basis for understanding the effects of published mutagenesis and biochemical modifications on the toxic activity of pneumolysin. In addition, spectroscopic data are presented that shed new light on pneumolysin activity and have guided us to hypothesise a detailed model of membrane insertion. These data show that the environment of some tryptophan residues changes on insertion and/or pore formation. In particular, spectroscopic analysis of a tryptophan mutant, W433F, suggests it is the residue mainly responsible for the observed effects. Furthermore, there is no change in the secondary structure content when the toxin inserts into membranes. Finally, the basis of the very low activity shown by a pneumolysin molecule from another strain of S. pneumoniae may be due to the movements of a key domain-domain interface. The molecular basis of pneumolysin-induced complement activation may be related to the structural similarity of one of the domains of pneumolysin to Fc, rather than the presumed homology of the toxin to C-reactive protein as previously suggested. Copyright 1998 Academic Press.

ANSWER 3 OF 27 MEDLINE

DUPLICATE 3

308-4994

ACCESSION NUMBER:

1998323373

MEDLINE

DOCUMENT NUMBER:

98323373

TITLE:

Application of molecular typing to the epidemiology

of Streptococcus pneumoniae.

AUTHOR:

Hall L M

CORPORATE SOURCE:

Department of Medical Microbiology, St Bartholomew's,

London, UK.. 1.m.c.hall@mds.qmw.ac.uk

SOURCE:

JOURNAL OF CLINICAL PATHOLOGY, (1998 Apr) 51 (4)

270-4. Ref: 45

Journal code: HT3. ISSN: 0021-9746.

PUB. COUNTRY:

ENGLAND: United Kingdom

Journal; Article; (JOURNAL ARTICLE)

General Review; (REVIEW)

(REVIEW, ACADEMIC)

LANGUAGE:

English

FILE SEGMENT:

Abridged Index Medicus Journals; Priority Journals;

Cancer Journals

ENTRY MONTH:

199809

ENTRY WEEK:

19980903

The spread of antibiotic resistance and the development of new AΒ vaccines have focused attention on the epidemiology of Streptococcus pneumoniae over recent years. While serotyping and the determination of antibiotic resistance remain primary methods for characterising pneumococci, molecular typing can add greater discrimination and complementary information. Methods based on restriction fragment length polymorphism within total DNA or non-specific

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polymerase chain reaction provide information representative of the whole genome and can be used to recognise closely related isolates from different sources, whether in the investigation of possible cross infection at the local level or in the investigation of national or international spread of antibiotic resistant strains. Fingerprinting of penicillin binding protein genes adds further information in the analysis of penicillin resistant isolates. The use of a combination of typing methods to analyse both the genome as a whole and specific loci has led to the realisation that pneumococci undergo horizontal gene transfer much more often than most other bacterial species. In particular the spread of penicillin resistance has been characterised by a combination of the spread of epidemic strains, transfer of chromosomal resistance genes from such strains into other genetic backgrounds, and transfer of capsule genes resulting in the switch of serotypes within strains. In the future molecular typing will have an important role in discovering whether widespread vaccination leads to genetic modification of the pneumococcal population causing invasive disease.

L5 ANSWER 4 OF 27 BIOSIS COPYRIGHT 1999 BIOSIS DUPLICATE 4

ACCESSION NUMBER: 1997:486725 BIOSIS DOCUMENT NUMBER: PREV199799785928

TITLE: Dissemination of antibiotic resistance.

AUTHOR(S): Roy, Paul H.

CORPORATE SOURCE: Cent. Rech. Cent. Hosp. Univ. Laval, Dep. Biochim.,

Univ. Laval, 2705 boulevard Laurier, Sainte-Foy, PQ

G1V 4G2 Canada

SOURCE: M-S (Medecine Sciences), (1997) Vol. 13, No. 8-9, pp.

927-933.

ISSN: 0767-0974. General Review

LANGUAGE: French

DOCUMENT TYPE:

SUMMARY LANGUAGE: French; English

While antibiotics have, for the past fifty years, been "miracle drugs", we are presently facing the "end of the miracle". The increasing use of antibiotics has led to the selection of bacteria resistant to multiple antibiotics. Diverse mechanisms of resistance are found in resistant bacteria. Among these are enzymatic degradation or alteration of antibiotic molecules (e.g. beta-lactamases and aminoglycoside modifying enzymes), altered targets (e.g. penicillin-binding proteins and dihydrofolate reductase), and drug efflux (e.g. of tetracycline). Often point mutations can drastically alter the enzyme or the target: beta-lactamases become able to digest third-generation cephalosporins, dihydrofolate reductase becomes resistant to trimethoprim, and DNA gyrase becomes resistant to quinolones. Resistance genes have not always been present in common pathogenic bacteria, but have been evolving in antibiotic producing bacteria or Searcher Shears 308-4994

in those cohabiting with them in the environment, and have recently been acquired by horizontal transfer. Many resistance genes are on conjugative plasmids of wide host range, often as part of transposons. Examples are the TEM beta-lactamase, whose gene can mutate to yield resistance to third-generation cephalosporins, and vancomycin resistance in enterococci, where a complete metabolic pathway for an altered cell wall in encoded by a transposon. In addition, a novel DNA element called an integron has been described, in which individual resistance genes exist as mobile cassettes and are rearranged by site-specific recombination, in a sort of natural genetic engineering, to form strongly expressed multiresistance operons. Knowledge of the mechanisms of resistance gene evolution and dissemination and of antibiotic usage patterns leads to the prediction, in a more or less immediate future, of the emergence of vancomycin-resistant staphylococci, of multiresistant pneumococci, and of third-generation-cephalosporin-resistant Haemophilus and Neisseria, for which the medical community must be prepared.

L5 ANSWER 5 OF 27 MEDLINE DUPLICATE 5

ACCESSION NUMBER: 96254545 MEDLINE

DOCUMENT NUMBER: 96254545

TITLE: In vitro activities of U-100592 and U-100766, novel

oxazolidinone antibacterial agents.

AUTHOR: Zurenko G E; Yagi B H; Schaadt R D; Allison J W;

Kilburn J O; Glickman S E; Hutchinson D K; Barbachyn

M R: Brickner S J

CORPORATE SOURCE: Pharmacia & Upjohn, Inc., Kalamazoo, Michigan 49001,

USA.

SOURCE: ANTIMICROBIAL AGENTS AND CHEMOTHERAPY, (1996 Apr) 40

(4) 839-45.

Journal code: 6HK. ISSN: 0066-4804.

PUB. COUNTRY: United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 199612

AB Oxazolidinones make up a relatively new class of antimicrobial agents which possess a unique mechanism of bacterial protein synthesis inhibition. U-100592 {(S)-N-[[3-[3-fluoro-4-[4-(hydroxyacetyl)-1-piperazinyl]-phenyl]-2-oxo-5-oxazolidinyl]methyl]-acetamide} and U-100766 {(S)-N-[[3-[3-fluoro-4-(4-morpholinyl)phenyl]-2-oxo-5-oxazolidinyl]methyl]-acetamide} are novel oxazolidinone analogs from a directed chemical modification program. MICs were determined for a variety of bacterial clinical isolates; the respective MICs of U-100592 and U-100766 at which 90% of isolates are inhibited were as follows: methicillin-susceptible Staphylococcus aureus, 4 and 4 micrograms/ml; methicillin-resistant S. aureus, 4 and 4

micrograms/ml; methicillin-susceptible Staphylococcus epidermidis, 2 and 2 micrograms/ml; methicillin-resistant S. epidermidis, 1 and 2 micrograms/ml; Enterococcus faecalis, 2 and 4 micrograms/ml; Enterococcus faecium, 2 and 4 micrograms/ml; Streptococcus pyogenes, 1 and 2 micrograms/ml; Streptococcus pneumoniae, 0.50 and 1 microgram/ml; Corynebacterium spp., 0.50 and 0.50 micrograms/ml; Moraxella catarrhalis, 4 and 4 micrograms/ml; Listeria monocytogenes, 8 and 2 micrograms/ml; and Bacteroides fragilis, 16 and 4 micrograms/ml. Most strains of Mycobacterium tuberculosis and the gram-positive anaerobes were inhibited in the range of 0.50 to 2 micrograms/ml. Enterococcal strains resistant to vancomycin (VanA, VanB, and VanC resistance phenotypes), pneumococcal strains resistant to penicillin, and M. tuberculosis strains resistant to common antitubercular agents (isoniazid, streptomycin, rifampin, ethionamide, and ethambutol) were not cross-resistant to the oxazolidinones. The presence of 10, 20, and 40% pooled human serum did not affect the antibacterial activities of the oxazolidinones. Time-kill studies demonstrated a bacteriostatic effect of the analogs against staphylococci and enterococci but a bactericidal effect against streptococci. The spontaneous mutation frequencies of S. aureus ATCC 29213 were <3.8 x 10(-10) and $<8 \times 10(-11)$ for U-100592 and U-100766, respectively. Serial transfer of three staphylococcal and two enterococcal strains on drug gradient plates produced no evidence of rapid resistance development. Thus, these new oxazolidinone analogs demonstrated in vitro antibacterial activities against a variety of clinically important human pathogens.

L5 ANSWER 6 OF 27 EMBASE COPYRIGHT 1999 ELSEVIER SCI. B.V.

ACCESSION NUMBER: 96071271 EMBASE

DOCUMENT NUMBER: 1996071271

TITLE: Bacterial resistance to carbapenems.

AUTHOR: Livermore D.M.

CORPORATE SOURCE: Department of Medical Microbiology, London Hospital

Medical College, Turner Street, London El 2AD, United

Kingdom

SOURCE: Advances in Experimental Medicine and Biology, (1996)

390/- (25-48).

ISSN: 0065-2598 CODEN: AEMBAP

COUNTRY: United States

DOCUMENT TYPE: Journal; Conference Article

English

FILE SEGMENT: 004 Microbiology

037 Drug Literature Index

LANGUAGE: English

SUMMARY LANGUAGE:

AB The carbapenems have the broadest spectra of all .beta.-lactams but resistance still occurs, caused by target modification, impermeability or .beta.-lactamase production. Target

modification or replacement is important in methicillin-Searcher : Shears 308-4994

resistant staphylococci, E. faecium and some pneumococci. These organisms present the greatest current threat to carbapenem efficacy. Impermeability to carbapenems arises in P. aeruginosa mutants, where it is contingent on loss of D2 outer membrane protein, a minor porin. This resistance functions only if the Pseudomonas retains its chromosomal group 1 .beta.-lactamase, and so reflects the interplay of impermeability and hydrolysis rather than impermeability alone; nevertheless porin loss is the critical change that engenders resistance. Resistance based on impermeability coupled to a group 1 .beta.-lactamase has also been described in E. cloacae, but demands loss of a major porin and is much rarer and less stable than in P. aeruginosa. Although group 1 .beta.-lactamases contribute to resistance in these organisms, they have only feeble carbapenemase activity. Chromosomal .beta.-lactamases with potent carbapenemase activity occur in most or all X. maltophilia. A. hydrophila and F. odoratum isolates. These enzymes, which cause carbapenem resistance when expressed copiously, are all zinc-dependent. Zinc carbapenemases also are a concern in B. fragilis, where they are encoded by the chromosomal DNA of c. 3% of isolates, though expressed by only 1%. Carbapenemases are extremely rare outside these species. Nevertheless, a plasmidic zinc carbapenemase was reported from one P. aeruginosa isolate and from several S. marcescens. Further carbapenemases, some not zinc-dependent, are known from a tiny numbers of Serratia, Enterobacter, and Acinetobacter isolates. Despite these various modes of resistance, carbapenems have retained their efficacy far heller than have expanded-spectrum cephalosporins. Whether this advantage will be retained indefinitely is uncertain. If resistance does become more prevalent it may be possible to derivatize the carbapenems so as to extend their activity. There is already interest in the design of carbapenems that bind .beta.-lactamresistant PBPs and, to an extent, in the development of carbapenemase inhibitors.

L5 ANSWER 7 OF 27 MEDLINE

ACCESSION NUMBER: 96342056 MEDLINE

DOCUMENT NUMBER: 96342056

TITLE: Bacterial resistance to carbapenems.

AUTHOR: Livermore D M

CORPORATE SOURCE: Department of Medical Microbiology, London Hospital

Medical College, UK.

SOURCE: ADVANCES IN EXPERIMENTAL MEDICINE AND BIOLOGY, (1995)

390 25-47. Ref: 80

Journal code: 2LU. ISSN: 0065-2598.

PUB. COUNTRY: United States

Journal; Article; (JOURNAL ARTICLE)

General Review; (REVIEW)

(REVIEW, TUTORIAL)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 199701 ENTRY WEEK: 19970104

AΒ

The carbapenems have the broadest spectra of all beta-lactams but resistance still occurs, caused by target modification, impermeability or beta-lactamase production. Target modification or replacement is important in methicillin-resistant staphylococci, E. faecium and some pneumococci. These organisms present the greatest current threat to carbapenem efficacy. Impermeability to carbapenems arises in P. aeruginosa mutants, where it is contingent on loss of D2 outer membrane protein, a minor porin. This resistance functions only if the Pseudomonas retains its chromosomal group 1 beta-lactamase, and so reflects the interplay of impermeability and hydrolysis rather than impermeability alone; nevertheless porin loss is the critical change that engenders resistance. Resistance based on impermeability coupled to a group 1 beta-lactamase has also been described in E. cloacae, but demands loss of a major porin and is much rarer and less stable than in P. aeruginosa. Although group 1 beta-lactamases contribute to resistance in these organisms, they have only feeble carbapenemase activity. Chromosomal beta-lactamases with potent carbapenemase activity occur in most or all X. maltophilia, A. hydrophila and F. odoratum isolates. These enzymes, which cause carbapenem resistance when expressed copiously, are all zinc-dependent. Zinc carbapenemases also are a concern in B. fragilis, where they are encoded by the chromosomal DNA of c. 3% of isolates, though expressed by only 1%. Carbapenemases are extremely rare outside these species. Nevertheless, a plasmidic zinc carbapenemase was reported from one P. aeruginosa isolate and from several S. marcescens. Further carbapenemases, some not zinc-dependent, are known from a tiny numbers of Serratia, Enterobacter, and Acinetobacter isolates. Despite these various modes of resistance, carbapenems have retained their efficacy far better than have expanded-spectrum cephalosporins. Whether this advantage will be retained indefinitely is uncertain. If resistance does become more prevalent it may be possible to derivatize the carbapenems so as to extend their activity. There is already interest in the design of carbapenems that bind beta-lactamresistant PBPs and, to an extent, in the development of carbapenemase inhibitors.

L5 ANSWER 8 OF 27 TOXLINE

ACCESSION NUMBER: 1994:52791 TOXLINE DOCUMENT NUMBER: CRISP-94-A04008-02

TITLE: SYNTHESIS OF OLIGONUCLEOTIDES.

AUTHOR: PROBST P G

CORPORATE SOURCE: FDA

U.S. DEPT. OF HEALTH AND HUMAN SERVICES; PUBLIC HEALTH SERVICE; NATIONAL INST. OF HEALTH, SYSOUT

CONTRACT NUMBER:

Z01BA04008-02

SOURCE:

(1992). Crisp Data Base National Institutes Of

Health. Award Type: G = Grant

DOCUMENT TYPE:

(RESEARCH)

FILE SEGMENT:

CRISP

LANGUAGE:

English

ENTRY MONTH:

199403

AB RPROJ/

RPROJ/CRISP Synthesis of oligonucleotides by our laboratory in collaborative efforts with several laboratories within the Division of Bacterial Products has proven invaluable to the following research programs: A PCR approach to clone and express a potential surface antigen from Mycoplasma pneumonia, and to modify existing cloning vectors. Development of a method to allow for direct cloning of PCR generated products, and as templates for DNA sequencing in Mycoplasma. Completion of the nucleotide sequence analysis of mycobacterial genes encoding several immunoreactive antigens. Isolation of the pneumococcal

pneumolysin gene from two types of genomic DNA, to produce trunicated point mutations of the pneumolynsin gene, and for conjugation of a single polysaccharide. Continued research in the production of selective mutation and amplification of bacterial toxin genes for use in examining the effect of mutation on toxin function. Primers were also utilized in the development of PCT protocols to rapidly detect non tuberculosos mycobacterial infections. Research in our laboratory is being conducted in the use of cloned overlapping strnads, to develop an in vitro transcription probe for allergenic screening of apple to determine and isolate the allergenic component. Studies are also continuing to include the use of oligonucleotides as tools for the creation of an epitope library.

L5 ANSWER 9 OF 27 MEDLINE

ACCESSION NUMBER:

93072694 MEDLINE

DOCUMENT NUMBER:

93072694

TITLE:

Modification of resistance to Streptococcus

pneumoniae by dietary ethanol, immunization, and

murine retroviral infection.

AUTHOR:

Darban H; Watson R R; Darban J R; Shahbazian L M

Department of Family and Community Medicine,

University Medical Center, Tucson, Arizona.

CONTRACT NUMBER:

CORPORATE SOURCE:

AA08037 (NIAAA)

SOURCE:

ALCOHOLISM, CLINICAL AND EXPERIMENTAL RESEARCH, (1992

Oct) 16 (5) 846-51.

Journal code: 35X. ISSN: 0145-6008.

PUB. COUNTRY:

United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Priority Journals

Searcher

Shears 308-4994

ENTRY MONTH:

199302

Hallmarks of the acquired immune deficiency syndrome (AIDS) are immunologic alterations, frequently associated with opportunistic infections. To study such associations, LP-BM5 murine retrovirus infection was used as a murine model of AIDS. Retrovirally infected and uninfected mice were fed a 5% (v/v) ethanol diet for 55 days and then fed a 7% v/v ethanol diet for the final 7 days to assert the role of ethanol as a cofactor in development of murine AIDS. There was a reduction in polymorphonuclear neutrophils count in ethanol-fed groups. Neutrophils increased in retrovirus-infected groups, except those vaccinated 10 days before challenge with live bacteria. The percentage of splenic lymphocytes in the retrovirus-infected group was reduced in comparison with controls. Survival of the mice challenged intraperitoneally with Streptococcus pneumoniae was increased by vaccination and suppressed by dietary alcohol. Retrovirus infection caused a much faster death rate after bacterial challenge than nonretrovirus infected controls. Vaccination played an important role in delaying the death rate in all treated groups. Transferring spleen cells from healthy, unimmunized mice also enabled the retrovirally infected mice to survive the bacterial infection longer. Enhancement of resistance to S. pneumoniae by vaccination and transfer of immunocompetent cells to mice immunosuppressed by retroviral infection show the potential to use immunomodulation to affect disease resistance in AIDS.

L5 ANSWER 10 OF 27 MEDLINE **DUPLICATE 6**

ACCESSION NUMBER:

92105031 MEDLINE

DOCUMENT NUMBER:

92105031

TITLE:

Truncated forms of PspA that are secreted from

Streptococcus pneumoniae and their use in functional

studies and cloning of the pspA gene.

AUTHOR:

Yother J; Handsome G L; Briles D E

CORPORATE SOURCE:

Department of Microbiology, University of Alabama,

Birmingham 35294..

CONTRACT NUMBER:

AI28457 (NIAID) AI21458 (NIAID) HD17812 (NICHD)

SOURCE:

JOURNAL OF BACTERIOLOGY, (1992 Jan) 174 (2) 610-8.

Journal code: HH3. ISSN: 0021-9193.

PUB. COUNTRY:

United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Priority Journals

ENTRY MONTH:

199204

AB Insertion-duplication mutagenesis was used to generate mutants of Streptococcus pneumoniae that produced truncated forms of PspA (pneumococcal surface protein A). The truncated products, representing from 20 to 80% of the complete PspA molecule, were all secreted from the cell and could be detected in Searcher : Shears 308-4994

unconcentrated culture medium. Analysis of the truncated molecules showed that the antigenic variability known to be associated with PspA is located in the alpha-helical N-terminal half of the molecule. This region was also found to contain immunogenic and protection-eliciting epitopes and to define the maximum region of the molecule that is likely to be surface exposed. The apparent molecular weight variability seen for PspA molecules of different S. pneumoniae strains was localized to both the N- and C-terminal halves of the protein. Attachment of PspA to S. pneumoniae was found to require regions located carboxy to the fifth repeat unit in the C-terminal end of the molecule. From the insertion-duplication mutants, the complete pspA gene was cloned and expressed in Escherichia coli. Differences in apparent molecular weight were observed when the same cloned product was expressed in E. coli and S. pneumoniae, suggesting that PspA is modified differently in the two hosts.

ANSWER 11 OF 27 MEDLINE

DUPLICATE 7

ACCESSION NUMBER:

92261297

MEDLINE

DOCUMENT NUMBER:

92261297

TITLE:

Comparative efficacy of autolysin and

pneumolysin as immunogens protecting mice

against infection by Streptococcus pneumoniae.

AUTHOR:

Lock R A; Hansman D; Paton J C

CORPORATE SOURCE:

Department of Microbiology, Adelaide Children's

Hospital, North Adelaide, South Australia.

SOURCE:

MICROBIAL PATHOGENESIS, (1992 Feb) 12 (2) 137-43.

Journal code: MIC. ISSN: 0882-4010.

PUB. COUNTRY:

ENGLAND: United Kingdom

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Priority Journals

ENTRY MONTH:

199208

Previous studies on Streptococcus pneumoniae have established that ΔR the pneumococcal proteins autolysin (N-acetylmuramyl-Lalanine amidase) and pneumolysin both contribute significantly to the virulence of the organism. In the present work, autolysin and a defined toxoid derivative of pneumolysin were tested, individually and in combination, for efficacy in a mouse model as antigens protecting against challenge with virulent, wild-type S. pneumoniae. While each antigen alone provided significant protection, the degree of protection was not increased when the antigens were administered together. In an additional experiment, mice were challenged with a genetically-modified mutant strain of pneumococcus unable to express active pneumolysin. Pre-immunization of such mice with autolysin failed to provide any significant protection against the challenge. The results of this study suggest that the most important contribution made by autolysin to the virulence of S. pneumoniae may Shears 308-4994 Searcher :

be its role in mediating the release of **pneumolysin** from the **pneumococcal** cytoplasm during infection.

L5 ANSWER 12 OF 27 MEDLINE

DUPLICATE 8

ACCESSION NUMBER:

92114766

DOCUMENT NUMBER:

92114766

TITLE:

Complement activation and antibody binding by

pneumolysin via a region of the toxin

MEDLINE

homologous to a human acute-phase protein.

AUTHOR:

Mitchell T J; Andrew P W; Saunders F K; Smith A N;

Boulnois G J

CORPORATE SOURCE:

Department of Microbiology, University of Leicester,

UK..

SOURCE:

MOLECULAR MICROBIOLOGY, (1991 Aug) 5 (8) 1883-8.

Journal code: MOM. ISSN: 0950-382X.

PUB. COUNTRY:

ENGLAND: United Kingdom

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Priority Journals

ENTRY MONTH:

199204

Pneumolysin, a membrane-damaging toxin, is known to activate the classical complement pathway. We have shown that 1 microgram ml-1 of pneumolysin can activate complement, which is a much lower level than observed previously. We have identified two distinct regions of pneumolysin which show homology with a contiguous sequence within acute-phase proteins, including human C-reactive protein (CRP). Site-directed mutagenesis of the pneumolysin gene was used to change residues common to pneumolysin and CRP. Some of the modified toxins had a reduced ability both to activate complement and bind antibody. We suggest that the ability of pneumolysin to activate complement is related to its ability to bind the Fc portion of immunoglobulin G.

L5 ANSWER 13 OF 27 MEDLINE

DUPLICATE 9

ACCESSION NUMBER:

92030189

DOCUMENT NUMBER:

92030189

TITLE:

Pneumolysin induces the salient histologic features of pneumococcal infection in the

rat lung in vivo.

AUTHOR:

Feldman C; Munro N C; Jeffery P K; Mitchell T J; Andrew P W; Boulnois G J; Guerreiro D; Rohde J A;

Todd H C; Cole P J; et al

CORPORATE SOURCE:

Department of Thoracic Medicine, National Heart and Lung Institute, Brompton Hospital, London, United

Kingdom.

SOURCE:

AMERICAN JOURNAL OF RESPIRATORY CELL AND MOLECULAR

BIOLOGY, (1991 Nov) 5 (5) 416-23. Journal code: AOB. ISSN: 1044-1549.

MEDLINE

PUB. COUNTRY:

United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Priority Journals

ENTRY MONTH:

199202

Streptococcus pneumoniae infections are common, but how they cause host tissue injury and death is incompletely understood. Immunization with pneumolysin, a thiol-activated toxin produced by the pneumococcus, partially protects animals during subsequent infection. The mechanism by which pneumolysin contributes to disease is not known. The aim of the present investigation was to determine the histologic changes induced by recombinant pneumolysin in the rat lung and to compare them with the changes induced by live organisms. Injection of either toxin (200 or 800 ng) or bacteria into the apical lobe bronchus was associated with the development of a severe lobar pneumonia restricted to the apical lobe. The changes induced by the toxin were greater at the higher concentration, and changes were most severe in those animals in which there was partial ligation of the apical lobe bronchus. The pneumonitis was less severe following injection of a modified toxin with decreased hemolytic

activity, generated by site-directed mutagenesis of the cloned pneumolysin gene, indicating that this property of the toxin was important in generating pulmonary inflammation. There was still considerable pneumonitis after injection of a modified toxin with decreased capacity to activate

MEDLINE

complement.

ANSWER 14 OF 27 MEDLINE

DUPLICATE 10

ACCESSION NUMBER:

89307578

DOCUMENT NUMBER:

89307578

TITLE:

Pneumolysin, the thiol-activated toxin of

Streptococcus pneumoniae, does not require a thiol

group for in vitro activity.

AUTHOR:

Saunders F K; Mitchell T J; Walker J A; Andrew P W;

Boulnois G J

CORPORATE SOURCE:

Department of Microbiology, University of Leicester,

United Kingdom..

SOURCE:

INFECTION AND IMMUNITY, (1989 Aug) 57 (8) 2547-52.

Journal code: GO7. ISSN: 0019-9567.

PUB. COUNTRY:

United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Priority Journals; Cancer Journals

:

ENTRY MONTH:

198910

The role of the single cysteine residue in the activity of the thiol-activated toxin pneumolysin was investigated using oligonucleotide-mediated, site-directed mutagenesis. Three modified toxins in which the cysteine residue was changed to

Searcher

Shears 308-4994

an alanine, a serine, or a glycine residue were purified to homogeneity and examined for activity. The Cys-428----Ala modified toxin was indistinguishable from the wild-type recombinant toxin in terms of hemolytic activity and lytic and inhibitory effects on human polymorphonuclear leukocytes (PMN), indicating that the cysteine residue is not essential for toxin activity. The Cys-428----Ser and Cys-429----Gly modified toxins had reduced activity on erythrocytes and polymorphonuclear leukocytes, being 6 and 20 times less active than the wild type, respectively. However, all the modified toxins formed oligomers in erythrocyte membranes to the same extent as the wild-type recombinant toxin. This suggests that the cysteine residue at position 428 is involved in neither the binding of toxin to membranes nor its insertion into the membrane, and also that the formation of oligomers is not by itself sufficient for toxin activity.

ANSWER 15 OF 27 MEDLINE **L5**

DUPLICATE 11

ACCESSION NUMBER:

89007040

MEDLINE

DOCUMENT NUMBER:

89007040

TITLE:

Stimulation of human neutrophils, monocytes, and

platelets by modified C-reactive protein (CRP) expressing a neoantigenic specificity.

AUTHOR:

Potempa L A; Zeller J M; Fiedel B A; Kinoshita C M;

Gewurz H

CORPORATE SOURCE:

Department of Immunology, Rush Presbyterian St.

Luke's Medical Center, Chicago, Illinois.

CONTRACT NUMBER:

R23AI23030 (NIAID)

HL-23457 (NHLBI) HL-00614 (NHLBI)

SOURCE:

INFLAMMATION, (1988 Aug) 12 (4) 391-405.

Journal code: GMO. ISSN: 0360-3997.

PUB. COUNTRY:

United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Priority Journals

ENTRY MONTH:

198901

AB C-reactive protein (CRP) can be structurally modified by heat, acid, or urea-chelation to express a neoantigen designated by us as neo-CRP. This antigen is also expressed on the in vitro primary protein translation products of both human and rabbit CRP. Unmodified CRP and CRP complexed with pneumococcal C-polysaccharide (CPS) do not express neo-CRP. Forms of CRP expressing neo-CRP but not native CRP antigenicity (even in the presence of CPS) consistently and in a dose-dependent manner potentiated the respiratory burst response of human polymorphonuclear leukocytes and peripheral blood monocytes to heat-modified IgG. Forms of CRP expressing neo-CRP antigenicity also induced reactions of aggregation and secretion

Shears 308-4994 Searcher :

from isolated platelets and potentiated platelet activation stimulated by ADP in platelet-rich-plasma, while native CRP alone or complexed with CPS again did not. Unlike CRP-CPS complexes, forms of CRP expressing neo-CRP were not able to activate the complement system. These data emphasize the biologic potential inherent in this humoral acute-phase reactant, particularly in the activation of the formed elements of the blood important in the inflammatory response. Since these cell-activating properties are preferentially observed when CRP is structurally modified to express the neo-CRP antigen, such a molecular conversion may be central to the structure-function relationships of CRP at local sites of inflammation and tissue injury.

L5 ANSWER 16 OF 27 MEDLINE DUPLICATE 12

ACCESSION NUMBER: 88105816 MEDLINE

DOCUMENT NUMBER: 88105816

TITLE: Overproduction and rapid purification of the amidase

of Streptococcus pneumoniae.

AUTHOR: Garcia J L; Garcia E; Lopez R

CORPORATE SOURCE: Centro de Investigaciones Biologicas, C.S.I.C.

Velazquez, Madrid, Spain..

SOURCE: ARCHIVES OF MICROBIOLOGY, (1987) 149 (1) 52-6.

Journal code: 7YN. ISSN: 0302-8933.

PUB. COUNTRY: GERMANY, WEST: Germany, Federal Republic of

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 198804

AB Oligonucleotide-directed mutagenesis of a plasmid containing the lytA gene coding for the pneumococcal amidase has allowed the separation of the coding sequence of the gene. This sequence has been placed in plasmid pIN-III(lppP-5)-A3 downstream from both a modified lipoprotein promoter and the lactose promoter to construct the recombinant plasmid pGL100. When Escherichia coli RB 791 (pGL100) was grown in the presence of lactose, the pneumococcal amidase accounted for 7% of the total protein present in this strain after 18 h incubation at 37 degrees C. The overproduced amidase was purified in a single-step procedure using a choline-Sepharose 6B column taking advantage of the fact that this enzyme was the unique protein with affinity for choline present in extracts obtained from E. coli RB791 (pGL100). The development of the above design opens up the possibility of studying the mechanism that regulates the activity of this important autolysin by using physiochemical techniques that require the availability of high amounts of purified amidase.

L5 ANSWER 17 OF 27 MEDLINE

DUPLICATE 13

ACCESSION NUMBER: 90074287 MEDLINE

DOCUMENT NUMBER: 90074287

TITLE: Gene conversion in Streptococcus pneumoniae.

AUTHOR: Sicard A M

CORPORATE SOURCE: Centre de Recherche de Biochimie et de Genetique

Cellulaires du C.N.R.S., Toulouse, France.. MICROBIOLOGIA, (1987 Feb) 3 (1) 5-12. Ref: 57

Journal code: AIF. ISSN: 0213-4101.

PUB. COUNTRY: Spain

SOURCE:

Journal; Article; (JOURNAL ARTICLE)

General Review; (REVIEW)

(REVIEW, TUTORIAL)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 199003

AB In pneumococcus, correction of mutations by the repair of mismatched bases results in gene conversion, i.e. transfer of genetic information from one strand of DNA to the other. Three different correction systems act upon a variety of mutations

: 1) A long-patch process (a few Kb) is efficient in the elimination of transitions, mostly +/- 1 base-pair mutations and less efficient in eliminating some transversions without affecting fairly long deletions. Neighboring sequences may interfere with this process. It is directed by at least two genes. 2) A localized conversion system acts on a six base-pair heteroduplex structure such as 5'ATTAAT/3'TAAGTA by specifically converting the mutated A base to the wild type C base.

Modifications of this configuration by site-directed mutagenesis lead to reduced conversion. 3) Fairly long deletions are eliminated during recombination by events that extend several scores of bases around the heterologus region. Although only the first conversion system has been shown to participate efficiently in protecting Streptococcus pneumoniae against spontaneous mutation, the two other processes may also eliminate mutations of different natures.

L5 ANSWER 18 OF 27 MEDLINE DUPLICATE 14

ACCESSION NUMBER: 85286297 MEDLINE

DOCUMENT NUMBER: 85286297

TITLE: Localized conversion in Streptococcus pneumoniae

recombination: heteroduplex preference.

AUTHOR: Sicard M; Lefevre J C; Mostachfi P; Gasc A M; Sarda C

SOURCE: GENETICS, (1985 Aug) 110 (4) 557-68.

Journal code: FNH. ISSN: 0016-6731.

PUB. COUNTRY: United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 198512

AB In pneumococcal transformation the frequency of recombinants between point mutations is generally

proportional to distance. We have recently described an aberrant marker in the amiA locus that appeared to enhance recombination frequency when crossed with any other allele of this gene. The hyperrecombination that we have observed in two-point crosses could be explained by two hypotheses: the aberrant marker induces frequent crossovers in its vicinity or the mutant is converted to wild type. In this report we present evidence showing that, in suitable three-point crosses, this hyperrecombination does not modify the recombination frequency between outside markers, suggesting that a conversion occurs at the site of this mutation. To estimate the length over which this event occurs, we isolated very closely linked markers and used them in two-point crosses. It appears that the conversion system removes only a few base pairs (from three to 27) around the aberrant marker. This conversion process is quite different from the mismatch-repair system controlled by hex genes in pneumococcus, which involves several thousand base pairs. Moreover, we have constructed artificial heteroduplexes using separated DNA strands. It appears that only one of the two heteroduplexes is specifically converted. The conversion system acts upon 5'..ATTAAT..3'/3'.. TAAGTA..5'. A possible role of the palindrome resulting from the mutation is discussed.

ANSWER 19 OF 27 MEDLINE

DUPLICATE 15

ACCESSION NUMBER:

85056632 MEDLINE

DOCUMENT NUMBER:

85056632

TITLE:

Opsonic activity of immunoglobulin prepared for

intravenous use.

AUTHOR:

Hetherington S V; Giebink G S

CONTRACT NUMBER:

AI-08821 (NIAID)

SOURCE:

JOURNAL OF LABORATORY AND CLINICAL MEDICINE, (1984

Dec) 104 (6) 977-86.

Journal code: IVR. ISSN: 0022-2143.

PUB. COUNTRY:

United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Abridged Index Medicus Journals; Priority Journals

ENTRY MONTH:

198503

The opsonic activity of two immunoglobulin preparations modified for intravenous infusion was tested against Streptococcus pneumoniae types 3, 7F, and 14 and two strains of Staphylococcus aureus by polymorphonuclear leukocyte uptake of 3H-thymidine-labeled bacteria. Reduced and alkylated immunoglobulin (Chem-IgG) and immunoglobulin prepared by chromatography with diethylaminoethyl-Sephadex (DEAE-IgG) were evaluated with and without complement and compared with the opsonic activity of immune serum globulin and heated pooled human serum. Opsonic activity of DEAE-IgG was greater than that of Chem-IgG and equivalent to the activity of immune serum globulin and pooled human

DUPLICATE 16

serum against S. aureus 502A and type 3 pneumococcus. Both intravenous immunoglobulins had lower opsonic activity than either pooled human serum or immune serum globulin against type 14 pneumococcus. There were no differences in antibody avidity for pneumococcal antigen among the immunoglobulins tested. All four opsonins had similar opsonic activity against the protein A-deficient S. aureus Wood 46. Modification of immunoglobulin for intravenous infusion by chemical alteration may adversely affect opsonic activity by changing the Fc portion of the antibody molecule.

L5 ANSWER 20 OF 27 MEDLINE

ACCESSION NUMBER: 84152108 MEDLINE

DOCUMENT NUMBER: 84152108

TITLE: Early pulmonary granulocyte recruitment in response

to Streptococcus pneumoniae.

AUTHOR: Vial W C; Toews G B; Pierce A K

CONTRACT NUMBER: HL-21817 (NHLBI)

SOURCE: AMERICAN REVIEW OF RESPIRATORY DISEASE, (1984 Jan)

129 (1) 87-91.

Journal code: 426. ISSN: 0003-0805.

PUB. COUNTRY: United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Abridged Index Medicus Journals; Priority Journals

ENTRY MONTH: 198406

Although polymorphonuclear leukocytes (PMN) are a conspicuous histologic feature of clinical and experimental pneumococcal pneumonia, neither the mechanism nor the magnitude of recruitment of these cells to the lung following lesser pneumococcal challenge is known. We have, therefore, investigated the early process of recruitment of PMN to alveolar spaces after pulmonary inoculation of Streptococcus pneumoniae in doses less than those causing pneumonia. We injected Balb/c mice with water and varying inoculums of pneumococci via an endobronchial catheter. Bronchoalveolar lavage (BAL) was performed on the inoculated lung at 0, 2, or 4 h after injection. Cellular response was measured and chemotactic activity was assayed on BAL supernatants at each time interval using the migration of human PMN through 3-micron filters in modified Boyden chambers by the leading front techniques. The BAL of normal and control animals (inoculum of sterile water only used for the control animals) yielded $5.03 +/- 1.51 \times 10(2)$ and $0.17 +/- 0.04 \times 10(5)$ PMN, respectively. The PMN recruitment at 4 h as a function of pneumococcal inoculum was described by the following equation: log PMN = 0.751 log Pn + 1.119 (r2 = 0.82, p less than 0.001). The PMN were, therefore, recruited in a dose-dependent manner. That recruitment may be caused by chemotactic substance(s) was suggested by the significant correlation between the PMN Shears 308-4994 Searcher

response and the distance of in vitro migration: $\log PMN = 0.057$ micron + 0.52 (r = 0.77, p less than 0.005). We have defined quantitatively the recruitment of PMN to the lung after pneumococcal challenge. (ABSTRACT TRUNCATED AT 250 WORDS)

L5 ANSWER 21 OF 27 MEDLINE

DUPLICATE 17

ACCESSION NUMBER:

84152107

MEDLINE

DOCUMENT NUMBER:

84152107

TITLE:

The role of C5 in polymorphonuclear

leukocyte recruitment in response to Streptococcus

pneumoniae.

AUTHOR:

Toews G B; Vial W C

SOURCE:

AMERICAN REVIEW OF RESPIRATORY DISEASE, (1984 Jan)

129 (1) 82-6.

Journal code: 426. ISSN: 0003-0805.

PUB. COUNTRY:

United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

AB

English

FILE SEGMENT:

Abridged Index Medicus Journals; Priority Journals

ENTRY MONTH:

198406

Polymorphonuclear leukocytes (PMN) play an important functional role in early pulmonary clearance of Streptococcus pneumoniae. The factors responsible for PMN recruitment to the lung after challenges with this organism are poorly defined. We used congenic C5-sufficient B10.D2/nSn (C5+) and C5-deficient B10.D2/oSn (C5-) mice to determine the importance of the C5 molecule in the PMN response to S. pneumoniae. The C5+ and C5- mice were injected with water and varying inoculums of pneumococci via an endobronchial catheter. Bronchoalveolar lavage (BAL) was performed on the inoculated lung at 0 and at 4 h after injection. Cellular response was measured and chemotactic activity was assayed in BAL supernatants at each time interval using human PMN in modified Boyden chambers by the leading front technique. Clearance of bacteria was studied by quantitative lung culture. The C5+ mice recruited significantly more PMN after challenges with both 10(4) and 10(6) pneumococci than did the C5- mice (p less than 0.05), but significant PMN accumulation did occur in C5- mice. Similarly, C5+ mice generated significantly more intraalveolar chemotactic activity than did C5- mice (p less than 0.05) but chemotactic activity was present in both C5+ and C5- mice in checkerboard assays. Pulmonary clearance of bacteria was significantly impaired in the absence of C5 at both inoculums (p less than 0.05). Our results indicate that the C5 molecule yields important PMN chemotaxins during the early time period after intrapulmonary inoculation of S. pneumoniae. However, PMN recruitment after this insult also results from other chemotaxins because both chemotactic activity and PMN recruitment occur within the alveoli of C5- mice.

DUPLICATE 18 ANSWER 22 OF 27 MEDLINE L5

83257856 MEDLINE ACCESSION NUMBER:

83257856 DOCUMENT NUMBER:

[Kinetics of the local immune response in lobar TITLE:

pneumonia].

Cinetique de la reponse immunitaire locale dans les

pneumonies lobaires.

Lafitte J J; Steenhouwer F; Roussel M; Courcol R; **AUTHOR:**

Voisin C; Caillaux M; Debosker Y; Geslin P

BULLETIN EUROPEEN DE PHYSIOPATHOLOGIE RESPIRATOIRE, SOURCE:

(1983 Mar-Apr) 19 (2) 209-13.

Journal code: BGX. ISSN: 0395-3890.

ENGLAND: United Kingdom PUB. COUNTRY:

Journal; Article; (JOURNAL ARTICLE)

French LANGUAGE:

Priority Journals FILE SEGMENT:

198311 ENTRY MONTH:

In 19 patients suffering from lobar pneumonia and treated with AB antibiotics, bronchoalveolar lavages were performed in attempt to follow the course of the biological disorders caused by the pulmonary bacterial infection. The cytologic study of the fluid harvested from 14 patients with normal immune reactivity showed, firstly, a polymorphonuclear leucocytosis and, after about ten days, a lymphocytosis, reaching 30 to 70% of the alveolar cell population. These cell disorders existed only in the lobe affected by the pneumonia process. In five alcoholics (one of them also splenectomized), the polymorphonuclear leucocytosis lasted 15 to 25 days and the lymphocytosis was delayed and moderate. We also searched for the pneumococcal antigen by counter-current immunoelectrophoresis using a polyvalent antiserum. We found it in 13 patients, 7 with a positive hemoculture for pneumococcus and 6 negatives. Clearance of this antigen was slow, non modified by alcoholism. We found this antigen in two patients later, between the 90th and 110th days, in the lavage fluid concentrated fifty times. The quantitative and qualitative study of the immunoglobulins revealed considerable individual variations, owing to the variable intensity of the local inflammation phenomenons and to the technical difficulties of their dosage in the lavage fluid.

DUPLICATE 19 ANSWER 23 OF 27 MEDLINE

MEDLINE ACCESSION NUMBER: 83141533

83141533 DOCUMENT NUMBER:

Mutagenesis in Streptococcus pneumoniae (TITLE:

pneumococcus) by transformation with DNA

modified by the carcinogen-mutagen,

aflatoxin B1.

Stark A A; Giroux C N AUTHOR:

MUTATION RESEARCH, (1983 Jan) 107 (1) 23-32. SOURCE:

Shears 308-4994 Searcher :

Journal code: NNA. ISSN: 0027-5107.

PUB. COUNTRY: Netherlands

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 198306

L5 ANSWER 24 OF 27 MEDLINE DUPLICATE 20

ACCESSION NUMBER: 76020435 MEDLINE

DOCUMENT NUMBER: 76020435

TITLE: A modifier mutation affecting

utilization of mannitol in pneumococcus.

AUTHOR: Rotheim M B; Hotchkiss R D

SOURCE: CANADIAN JOURNAL OF MICROBIOLOGY, (1975 Aug) 21 (8)

1139-43.

Journal code: CJ3. ISSN: 0008-4166.

PUB. COUNTRY: Canada

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 197602

AB The altered growth of a pneumococcal mutant

containing marker M, which confers ability to utilize mannitol, and a modifier gene is reported. The modifier gene

is closely linked to the M gene and imposes a requirement for growth

in 0.4% glucose before growth in mannitol medium. The tentative position of the M gene of ${\bf pneumococcus}$ is ery-r str-r M

sul-rd.

L5 ANSWER 25 OF 27 MEDLINE DUPLICATE 21

ACCESSION NUMBER: 74054088 MEDLINE

DOCUMENT NUMBER: 74054088

TITLE: Integration efficiency in DNA-induced transformation

of Pneumococcus. I. A method of

transformation in solid medium and its use for

isolation of transformation-deficient and

recombination-modified mutants.

AUTHOR: Tiraby G; Claverys J P; Sicard M A SOURCE: GENETICS, (1973 Sep) 75 (1) 23-33.

Journal code: FNH. ISSN: 0016-6731.

PUB. COUNTRY: United States

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English ENTRY MONTH: 197403

L5 ANSWER 26 OF 27 TOXLINE

ACCESSION NUMBER: 1991:40582 TOXLINE

DOCUMENT NUMBER: EMIC-16357

TITLE: INTEGRATION EFFICIENCY IN DNA-INDUCED TRANSFORMATION

OF PNEUMOCOCCUS. 1.A METHOD OF

TRANSFORMATION IN SOLID MEDIUM AND ITS USE FOR

ISOLATION OF TRANSFORMATION DEFICIENT AND

RECOMBINATION MODIFIED MUTANTS.

AUTHOR:

TIRABY G; CLAVERYS J; SICARD M A

MEDLINE

SOURCE:

GENETICS, (1973). Vol. 75, pp. 23-33.

CODEN: GENTAE.

DOCUMENT TYPE:

(ORIGINAL DATA)

FILE SEGMENT:

EMIC

LANGUAGE:

Unavailable

ENTRY MONTH:

199103

ANSWER 27 OF 27 MEDLINE **L**.5

ACCESSION NUMBER:

69161419

DOCUMENT NUMBER: TITLE:

69161419 [Bacteria modified by parietal alteration.

Streptococci, staphylococci, pneumococci; their frequency in hemocultures. Attempted

interpretation).

Les bacteries modifiees par alteration de la paroi. Streptocoques, Staphylocoques,

Pneumocoques; leur frequence dans les hemocultures.

Essai d'interpretation.

AUTHOR:

Nativelle R; Deparis M

SOURCE:

PRESSE MEDICALE, (1968 Nov 27) 76 (46) 2199-200.

Journal code: PLP. ISSN: 0032-7867.

PUB. COUNTRY:

France

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

French

FILE SEGMENT:

Priority Journals

ENTRY MONTH:

196907

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(FILE 'CAPLUS' ENTERED AT 16:07:05 ON 02 SEP 1999)

L6

1 SEA ABB=ON PLU=ON PNVJ1 OR PNVJ20 OR PNVJ22 OR PNVJ45 OR PNVJ56 OR PNV103 OR PNV207 OR PNV111 OR PNV21 OR PNVJ(W) (1 OR 20 OR 22 OR 45 OR 56) OR PNV(W) (103 OR 207 Named poly-peptides claims 7-15

OR 111 OR 211)

L7

O SEA ABB=ON PLU=ON L6 NOT L3

FILE 'MEDLINE, BIOSIS, EMBASE, JICST-EPLUS, TOXLINE, TOXLIT, CONFSCI, PHIC, PHIN' ENTERED AT 16:09:16 ON 02 SEP 1999

O SEA ABB=ON PLU=ON L6 L8

FILE 'HOME' ENTERED AT 16:10:11 ON 02 SEP 1999

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308-4994 Searcher : Shears